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ESULT 4
ABU52249
ID ABU52249 standard; protein; 119 AA.
XX
AC ABU52249;
XX
DT 07-MAY-2003 (first entry)
XX
DE Helicobacter pylori selected interacting domain (SID) protein #1593.
XX
KW Protein-protein interaction; ulcer; selected interacting domain; SID.
XX
OS Helicobacter pylori.
XX
PN WO200266501-A2.
XX
PD 29-AUG-2002.
XX
PF 28-DEC-2001; 2001WO-EP015428.
XX
PR 02-JAN-2001; 2001US-0259302P.
XX
PA (HYBR-) HYBRIGENICS.
PA (INSP) INST PASTEUR.
XX
PI Legrain P, Rain J, Colland F, De Reuse H, Labigne A;
XX
DR WPI; 2002-674910/72.
DR N-PSDB; ABX66994.
XX
PT New complexes of protein-protein interactions in Helicobacter pylori,
PT useful for identifying modulating compounds for treating or preventing
PT ulcers in mammals.
XX
PS Claim 6; Page 477; 642pp; English.
XX
CC The invention describes a complex of protein-protein interactions in
CC Helicobacter pylori selected from 421 complexes given in the
CC specification. The complex of protein-protein interactions are useful for
CC screening for agents which modulate the interaction of proteins.
CC Modulating compounds which binds to a targeted bacterial protein may be
CC used for treating or preventing ulcers in a human or animal. This is the
CC amino acid sequence of a selected interacting domain (SID), identified
CC via protein-protein interactions. Note: Where the patent number printed
CC at the top of the pages in the specification has obscured areas of
CC protein sequence, the indexer has replaced the residue with an X to
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CC represent an illegible residue

XX

SQ Sequence 119 AA;

Query Match 40.5%; Score 589; DB 5; Length 119; out of 285 for full length protein

Best Local Similarity 100.0%; Pred. No. 2.3e-36;

Matches 119; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 150

EAKEKSAPKQVTTKDIVKEKDKQEESNKTSEGATSEAQAYNPGVSNEFLMKIQT AISSKN 209

Db 1

EAKEKSAPKQVTTKDIVKEKDKQEESNKTSEGATSEAQAYNPGVSNEFLMKIQT AISSKN 60

Qy 210

RYPKMAQIRGIEGEVLVSFTINADGSVTDIKVVKSNTTDILNHAALEAIKSAAHLF PKP 268

Db 61

RYPKMAQIRGIEGEVLVSFTINADGSVTDIKVVKSNTTDILNHAALEAIKSAAHLF PKP 119